

iGEM 2016: Pittsburgh  
**Week 10 Lab Notebook**

*Monday, July 25*

Lab meeting

Phosphorylate PT3-RBS and PT3 (Maddie/Maya)

Resuspend to 100uM with water

[Phosphorylation](#)

Made [annealing buffer](#) (Maya)

[Liquid cultures](#) of Collins switches (Claire)

Fundraising emails (Claire)

*Tuesday, July 26*

[Miniprep](#) liquid cultures (Claire)

Concentrations:      D 1: 48.3      G 1: 72.5  
                                 D 2: 60.4      G 2: 63.1

[Cell-free](#) dilution time course with Collins plasmids (Claire)

D:	Solution A	2 uL		22 uL
	Solution B	1.5 uL		16.5 uL
	RNAse inhibitor	0.25 uL	x 11 =	2.75 uL
	Substrate	0.25 uL		2.75 uL
	Switch:	0.82 uL at 60.4 ng/uL		9.02 uL
	Trigger:	0.18 uL at 5 uM	x 8 =	1.44 uL
G:	Solution A	2 uL		20 uL
	Solution B	1.5 uL		15 uL
	RNAse inhibitor	0.25 uL	x 10 =	2.5 uL
	Substrate	0.25 uL		2.5 uL
	Switch:	0.59 uL at 84.1 ng/uL		5.9 uL
	Trigger:	0.36 uL at 2.5 uM	x 8 =	2.52 uL
	Water	0.05 uL	x 7 =	0.35 uL

Master mix:      Solution A      42 uL  
                         Solution B      31.5 uL  
                         RNAse inhibitor 5.25 uL  
                         Substrate      5.25 uL

Separate out 40 uL for G

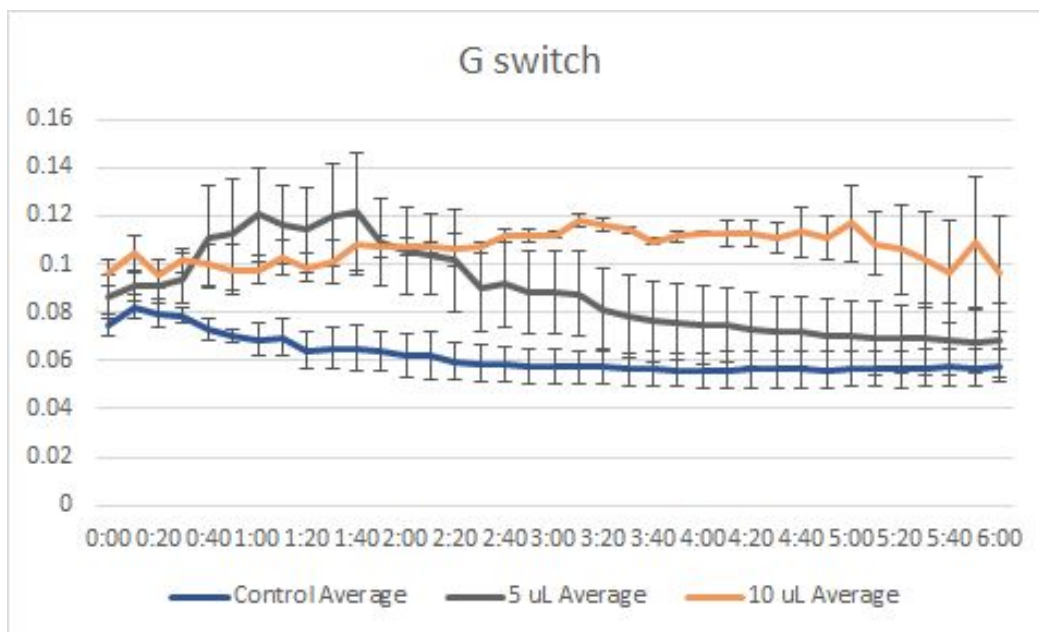
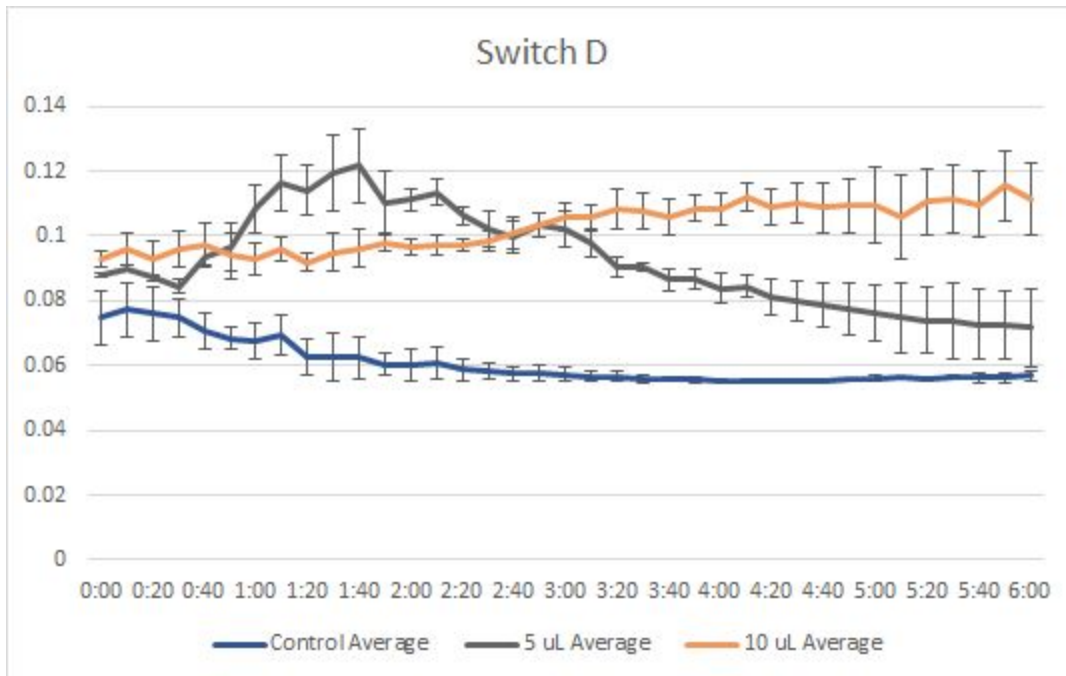
Add switches

Separate out 14.46 uL for D control //      13.77 uL for G control  
                         with 0.54 uL water      //      with 1.23 uL water

Add trigger to remaining

Add 5 uL to wells, dilute to appropriate volumes

6-hour time course with readings every minute, lid on



Dilute all wells to 10 uL to read final absorbances

Plate: Row K: D switch

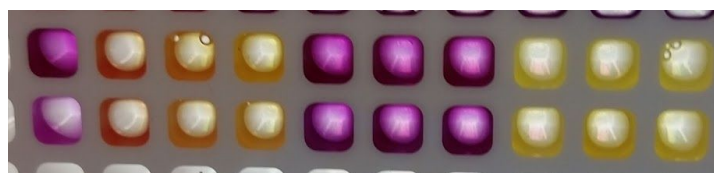
Columns 1-3: control

Column 10: leftover  
reaction incubated in  
heat block

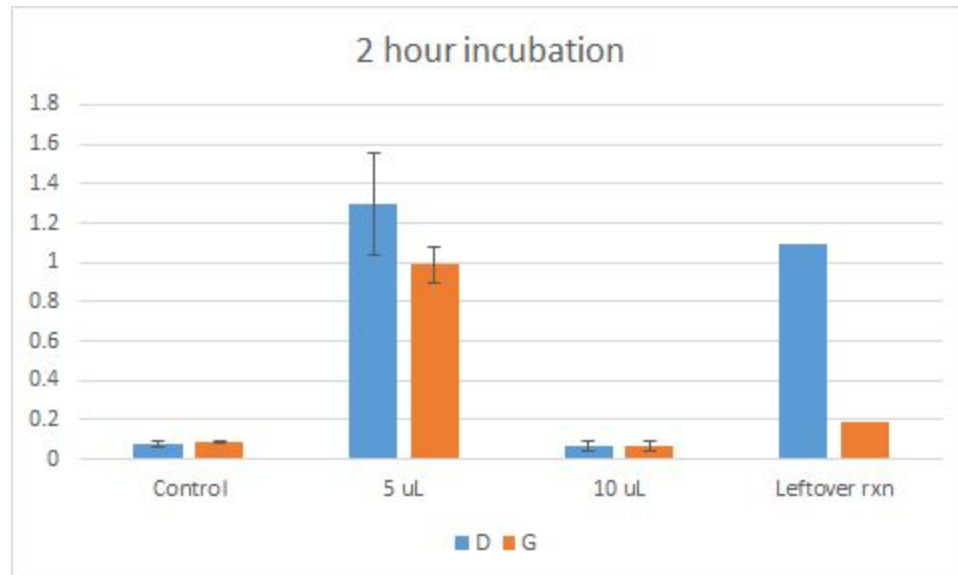
Row L: G switch

Columns 4-6: 5 uL

Columns 7-9: 10 uL



\*taken from bottom of plate



Results quite different from last week

5 uL reactions consistently proceeded

10 uL reactions barely orange

Put plate back in plate reader for overnight time course

[Anneal](#) (Maddie/Maya)

Sequences received

Was not LacZ like we believed

Found another file with LacZ in it, will transform that

Gel Extraction of RBS-T3 and Plasmid Backbone

[Digest](#)

5 ug DNA, 40 uL reaction

Cut with XbaI and PstI (2 uL each)

4 uL Fastdigest Buffer

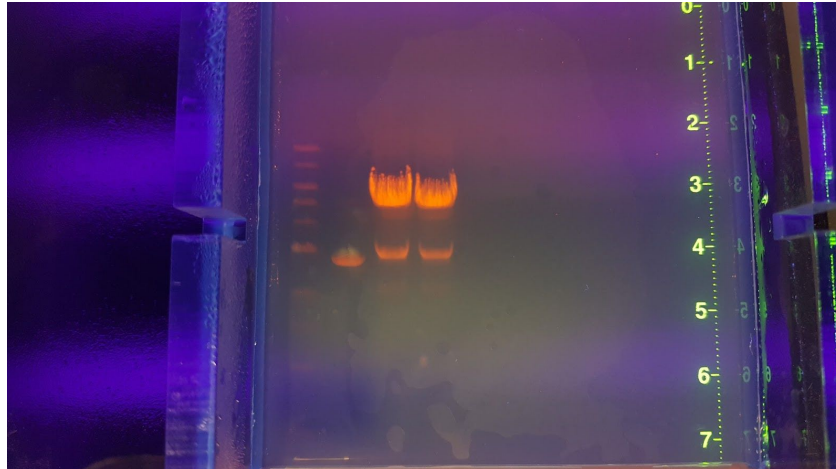
15 uL DNA (173.5 ng/uL)

8.-- uL DNA (--- ng/uL)

8.-- uL H2O

Gel Extract Digest Protocol (45 min at 37C, 20 min at 65C)

[Gel](#)



Lane 1: 1kb Ladder

Lane 2: Control

Lane 3: 20 uL Digest

Lane 4: 20 uL Digest

[Gel extract](#)

DNA concentration:

Fundraising (Claire)

Wednesday, July 27

[Cell-free](#) Collins dilution (Claire)

Overnight time course: wells dried up

10 uL reaction darker, but not fully purple



\*taken from bottom of plate

Time course data is function of evaporation more so than reaction

Modeling (Aife/Maya)

Talked with Natasa about modeling toehold and population/economics of lead

[Miniprep](#) Liquid Cultures (Maddie)

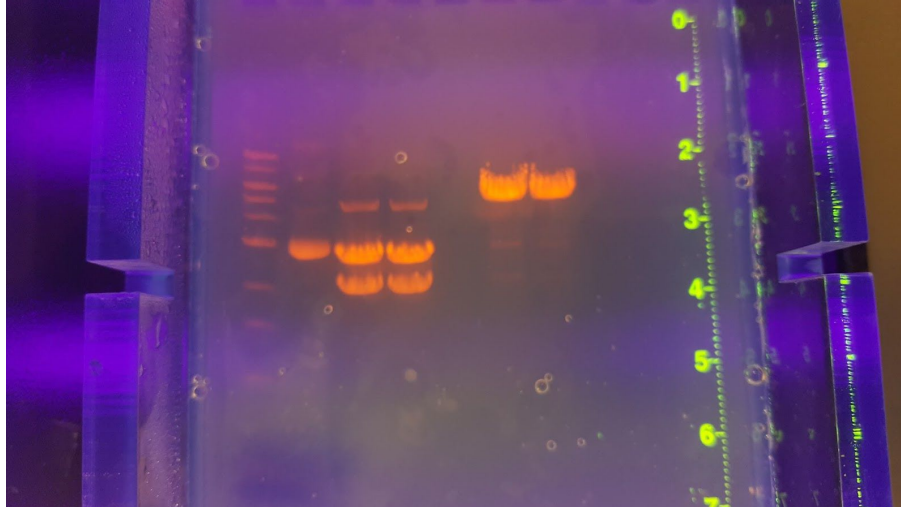
[Gel Extraction](#) of RBS-T3

[Digest](#) with EcoRI and PstI (Maddie)

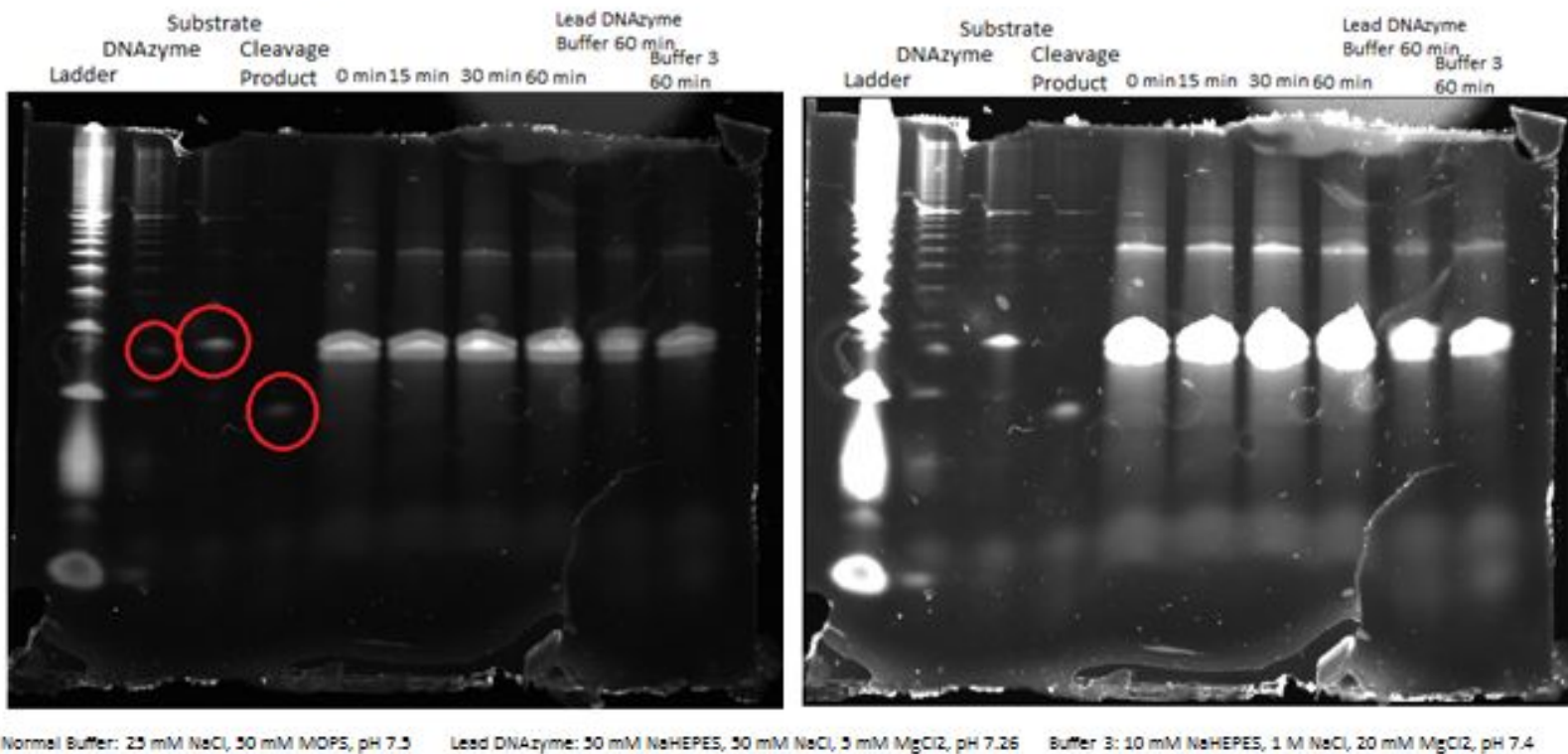
50 uL reaction

Incubate for 2 hours

[Gel](#)



Make CM [plates](#) (Maddie/Maya)  
[Liquid cultures](#) (Maya)  
 PT7-RBS-amiCP-Term  
 LacZ(?)  
[dPAGE](#) DNAzyme (Praneeth)



No evidence of cleavage

Thursday, July 28

[Camp BioE](#) presentation (Claire and Praneeth)

[Miniprep](#) Liquid Cultures (Maddie)

[Ligation](#) of PT3 and PT3-RBS with plasmid backbone ([Maya](#))

Control

3:1 PT3

7:1 PT3

3:1 PT3-RBS

7:1 PT3-RBS

[PCR Amplification](#) of T3 ([Maya](#))

At 55C and 52C, 20 uL reactions

*Friday, July 29th*

Prepare and ship LacZ and amilCP construct for [sequencing](#) (Claire)

LA01: LacZ VR

LA02: LacZ VF2

LA03: amilCP VR

LA04: amilCP VF2

15 uL reaction

5 uL 5 uM primer

500 ng plasmid = 1.26 uL amilCP // 5.44 uL lacZ

Water to volume = 8.74 uL // 5.44 uL

[Colony PCR](#) of PT3 and PT3-RBS ([Maya](#))

100 uL Master Mix ([http://2013.igem.org/Colony\\_PCR\\_Protocol](http://2013.igem.org/Colony_PCR_Protocol))

20 uL 5X Reaction Buffer

15 uL VR (5 uM)

15 uL VF2 (5 uM)

2 uL dNTP

2 uL DMSO

44 uL H2O

1 uL Phusion

10 uL per reaction

Used pipette tip to pick part of colony and dipped into tube

Used BioBrick Primer cycle in PCR machine

[Gel](#) (1.5%)

Lane 1: 1kb ladder

Lane 2: 10bp ladder

Lane 3: Plasmid Backbone

Lane 4: Colony 1

Lane 5: Colony 2

Lane 6: Colony 3

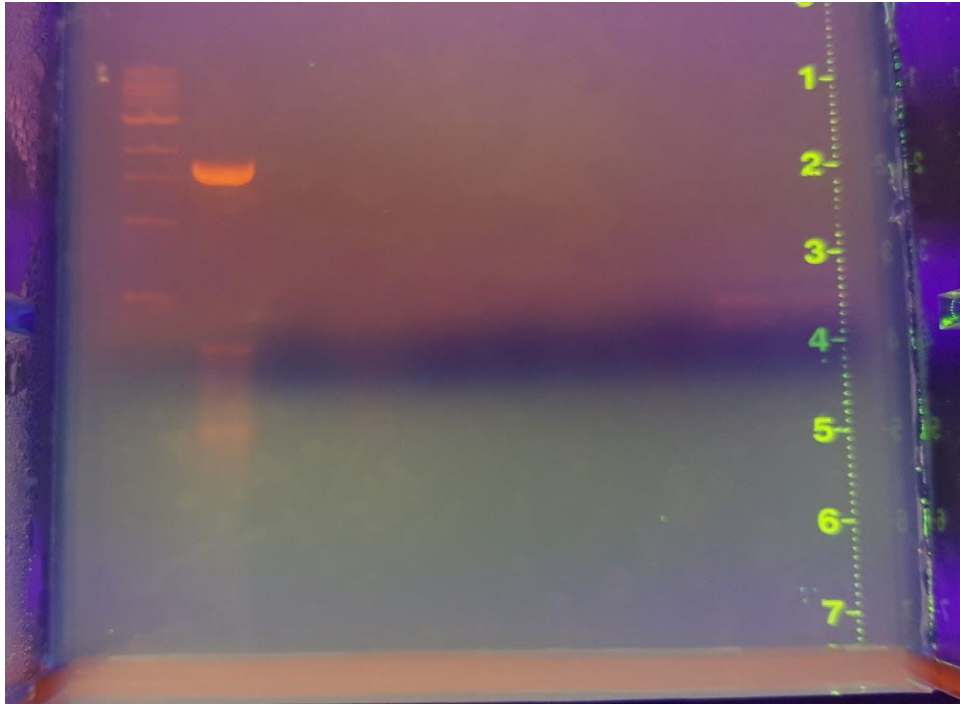
Lane 7: Colony 4

Lane 8: Colony 5

Lane 9: Colony 6

Lane 10: Colony 7





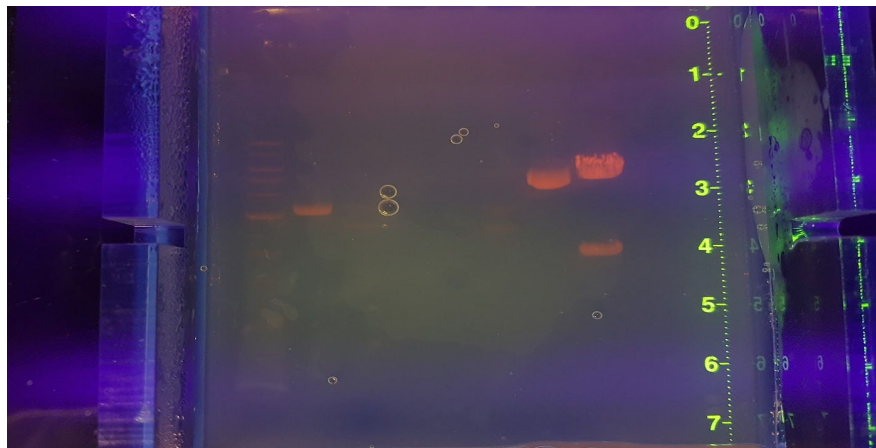
Not 100% sure what happened here:

Best guess is that control somehow (??) got into lane 2

Ran the gel too long

Use imager next time (Praneeth gone so no access)

T3 amplification check along with amilCP construct check (Maddie)



[PAGE](#) check annealing with Nick (Claire)

15% gel (poured in an extra mL of TBE)

16 uL sample + 4 uL loading buffer with bromophenol blue

2 uL 10 bp ladder with 1 uL loading buffer

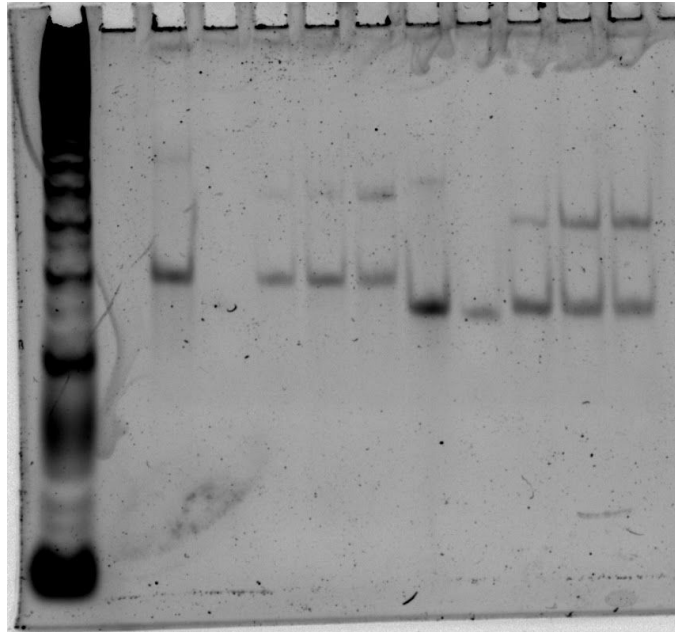
Lane	1	ladder	7	D 50 nM : 100 nM
	2	(empty)	8	G substrate, 50 nM
	3	D substrate, 50 nM	9	G catalytic, 50 nM
	4	D catalytic, 50 nM	10	G 50 nM : 50 nM

5 D 50 nM : 50 nM  
6 D 50 nM : 75 nM

11 G 50 nM : 75 nM  
12 G 50 nM : 100 nM

Run at 100V for 1 hour

SYBR Gold stain 20 min



Not sure what happened to D catalytic strand

Duplexes are formed (especially G 1:1.5 and G 1:2)

Excess substrate strand

Gel from 07-15 was just viewed from the bottom. Consistent with this gel

YAYYYYYY

Fundraising (Claire)